

Figurel

ATF1: QTTKTDDPQ-LKREIRLMKNREAARECR-RKKK (205 - 235 of SEQ ID NO:1)
CREB: PTQPAEE-AARKREVRLMKNREAARECR-RKKK (275 - 305 of SEQ ID NO:2)
GCN4: ----IDDPAALKR-AR---NTEAARRSRARKL- (224 - 247 of SEQ ID NO:3)

Figure 2

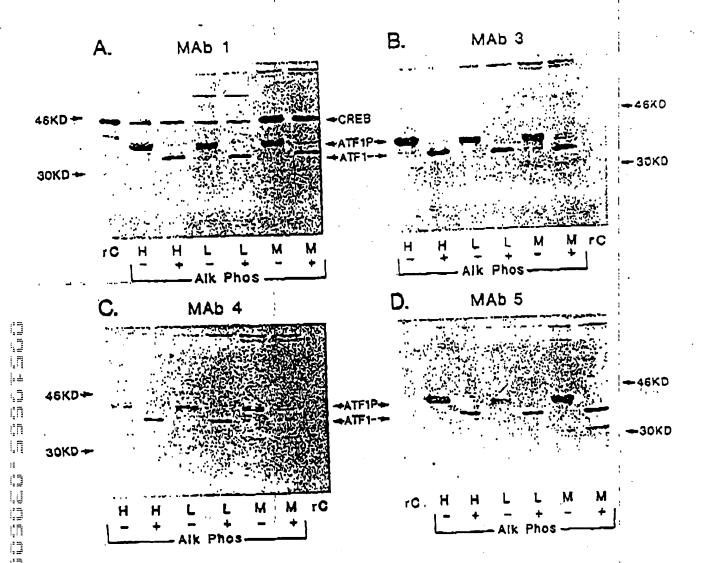
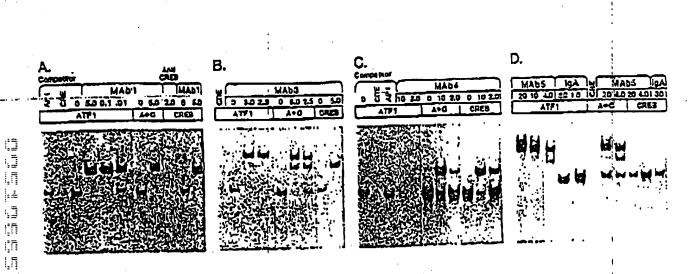


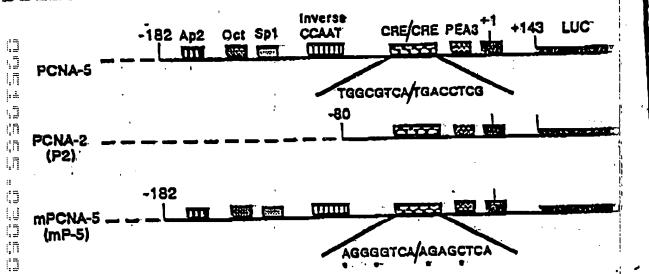
FIGURE 3

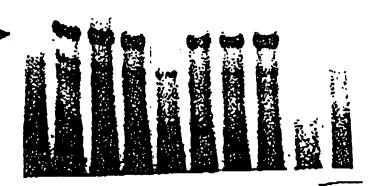


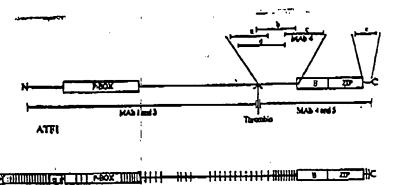
4... 4... ... 4... 4... 4... 4...

STATER A

PCNA-LUC constructs







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MAD1 MAD3 MAD4 MAD5

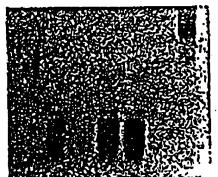
-46KD
-30KD
-30KD
-21.5KD

C-terminal
fragment
Thrombin 40 80 40 80 40 80 40 80

(n (n (n

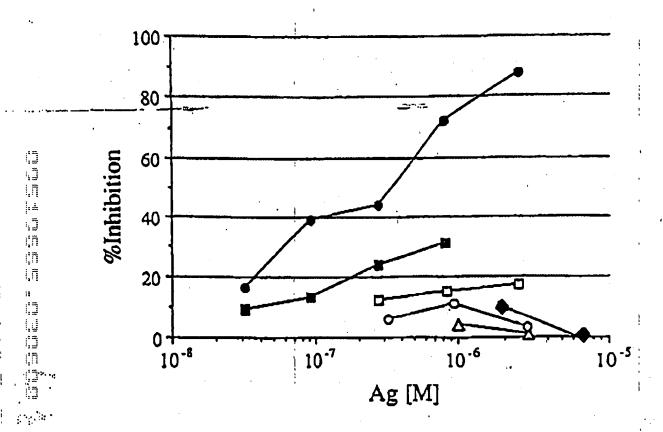
FIGURE 8

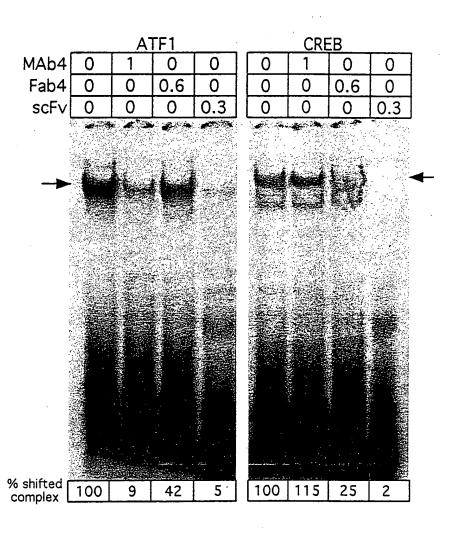
 $\mathcal{H}_{\mathbb{Q},d}^{\mathbb{Q}}$



M1 M3 M4 M5

FIGURE 9





Figurell

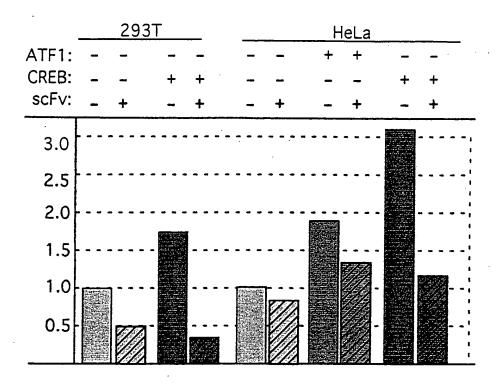


Figure 12

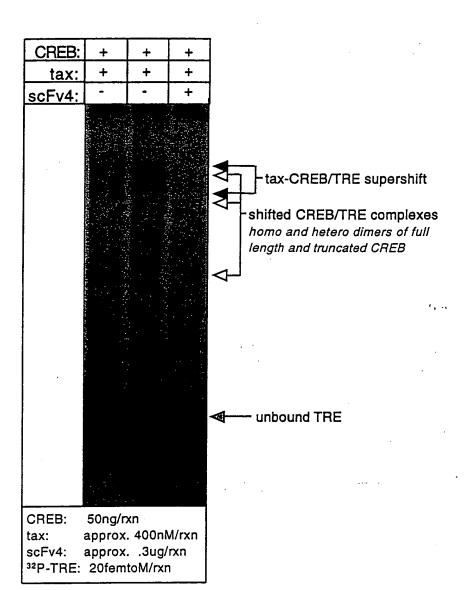


Figure 13